AMENDMENTS TO THE SPECIFICATION

In the specification:

Please replace the paragraph at page 5, lines 5-15, with the following amended paragraph:

Figure 2 shows the cDNA and deduced amino acid sequences of human *PIN1* and homologies with other WW domain proteins and PPIases. Figure 2A shows the predicted Pin1 amino acid sequence is indicated in one-letter code (SEQ ID NO:2). The fusion points between GAL4 and Pin1 in six different isolated clones were: clone H20 at C-9; clone H16, 24 and 38 at G+13; clones H7 and H36 at C+15. Underlined residues form a consensus bipartite nuclear localization signal. The N- and C-terminal boxes indicate the WW domain and PPIase domain. Nucleotide numbers are on the left and amino acid numbers on right (SEQ ID NOS:1 and 2, respectively). Figure 2B (SEQ ID NOS:8-15, respectively) and 2C (SEQ ID NOS:16-22, respectively) show alignments of the WW domain (B) and PPIase Domain (C) in selected proteins. Identical residues are shown in the bottom row. Dashes indicate gaps introduced to make the alignment. Cbf2, cell binding factor 2; SC, *S. cervisiae*; EC, *E. coli*; BS, *B. subtilis*; CJ, *C.jejuni*; AT, *A. thaliana*.